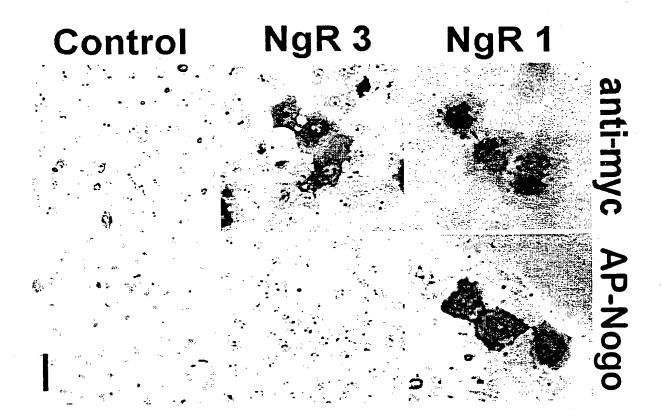
FIG. 1A

C	SCPMLCTCYS GCPRDCVCYP PCPGACVCYN -CPC-CY-	100 TFGSNLLT HFSPAMVT SFRACRNLTI -F	150 FQGLERLQSL FQGLVKLHAL FHGLGRLHTL F-GLLL	200 DLANLSHLFL DLVNLSHLFL DLGNLTHLFL DL-NL-HLFL	250 SRLTILYLFN HRLTTLFLFN GRLMTLYLFA -RLL-LF-
			150 RHLRSLEPDT FQGLERLQSL RQLRTLAPET FQGLVKLHAL AQLRSVDPAT FHGLGRLHTL LRP-T F-GLLL	LLHLQDDLFA IEYLQDDIFV LQALPDDTFR L-DD-F-	GVHRAAFRGL WVHHKAFHDL HVHPHAFRDL -VHAFL
	ACLLLML L SCCVELLLLL L AGGSRLLAWV L	NNFSSVPLSL PPSTQRLFLQ NNLIRTLRPG HNFAAIPEGI PEDSERIFLQ NNRITFLQQG QGLQAVPVGI PAASQRIFLH GNRISHVPAA		LQYLYLQENS LQYLYLQDNH LQYLYLQDNA LQYLYLQ-N-	
	LRRLLQAPAS ACLLLML QSPVQAAQVS GCCVELLLLL ~~~~MKRAS AGGSRLLAWV		IYPGTFRHLQ ALEELDLGDN IAPNTFEGFV HLEELDLGDN IDAAAFTGLA LLEQLDLSDN IFLE-LDL-DN	151 HLYRCQLSSL PCNIFRGLVS YLYKCGLSAL PAGIFGGLHS HLDRCGLQEL GPGLFRGLAA -LC-LLF-GL	201 HCNRLRLLTE HVFRGLGSLD HCNKLWSLGQ GIFRGLVNLD HCNRISSVPE RAFRGLHSLD HGNFRGLLD
	MSWQSGTTVT QSPVQAAQVS GCCVELLLL LAGELPLGG. MSWQSGTTVT QSPVQAAQVS GCCVELLLL LAGELPLGG. MSWQSGTTVT QSPVQAAQVS GCCVELLLL LAGELPLGG.	51 SP. PTVSCQA NNFSSVPLSL AP. MTVSCQA HNFAAIPEGI EPKVTTSCPQ QGLQAVPVGI -PT-SC	101 LWLFSNNLST LWIYSNNITF LWLHSNVLAR LWSN	151 HLYRCQESSL YLYKCGLSAL HLDRCGLQEL -LC-L	201 HCNRLRLLTE HCNKLWSLGQ HGNRISSVPE HGN
	· · · <u>-</u>			·	
	NOGO R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus	NOGO-R2 NOGO-R3 NOGO-R1 Consensus

Fig. 1B

300 NSLASLPGEA LADLPSLEFL RLNANPWACD CRARPLWAWF QRARVSSSDV NSLTELQGDC LAPLVALEFL RLNGNAWDCG CRARSLWEWL RRFRGSSSAV NNLSALPTEA LAPLRALQYL RLNDNPWVCD CRARPLWAWL QKFRGSSSEV N-LL LA-LL RLN-N-W-C- CRAR-LW-WR-SSS-V	350 TCATPPERQG RDLRALREAD FQACP.P AAPTRPGSRA PCATPELRQG QDLKLLRVED FRNCTGP VSPHQIKSHT PCSLPQRLAG RDLKRLAAND LQGCAVATGP YHPIWTGRAT DEEPLGLPKC -CPG -DLLDCPP	351 AOOSSSNH.LY G.VAE AGAPPADPS. TLYRDLPALTTSDRAA RKEHHPSH G. ASRDKGHP HGHPPGSRSG YKKAGKNCTS CQPDAADKAS VLEPGRPASA GNALKGRVPP GDSPPGNGSG PRHI.NDSPF	450 EDSRGR QCGDAPTE.D DYWCGY	451 ACQAPPDSRG PALSAGLPSP LLCLLLLVPH HL~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
NOGO-R2	NOGO - R2	NOGO-R2	NOGO-R2	NOGO-R2
NOGO-R3	NOGO - R3	NOGO-R3	NOGO-R3	NOGO-R3
NOGO-R1	NOGO - R1	NOGO-R1	NOGO-R1	NOGO-R1
Consensus	Consensus	Consensus	Consensus	Consensus

3/6 FIG. 2



4/6 FIG. 3

	-	5			50
Human NOGO-R1		MKRASA	GGSRLLAWVL	WLQAWQVAAP WLQAWRVATP	CPGACVCYNE
Murine NOGO-R1	ຜ		GCCVELLLL		CPRDCVCYPA
Human NOGO-R3 Human NOGO-R2	1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	GLRRLLQ			CPMLCTCYSS
Consensus	5 5 5 5 5 5	1 1 1 1 2 2 2 2 2	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	t	CPCY
	51 LRR NT	-		LRR 1	100
Human NOGO-R1	VTTSCPQQ		AASQRIFLHG		FRACRINLTIL
Murine NOGO-R1	PKVTTSCPQQ				FOSCRINITED AMVITE
Murine NOGO-R3	P.MTVSCQAH	P.MIVSCOAH NFAALFEGIF	VDSFRVFLON NRIGLLOPGH	NR I GLLOPGH	FSP. AMVTL
Human NOGO-R2	P. PTVSCQAN	P. PTVSCQAN NFSSVPLSLP	PSTQRLFLON NLIRTLRPGT	NLIRTLRPGT	FGSNLLTL
Consensus	PT-SC			N-IN	FI
	101 LRR 2	2	LR	LRR 3	150
Human NOGO-R1	WLHSNVLARI	DAAAFTGLAL	LEQLDLSDNA	QLRSVDPATF	
Murine NOGO-R1	WLHSNALARI	DAAAFTGLTL	LEQLDLSDNA	QLHVVDPTTF	HGLGHLHTLH
Murine NOGO-R3	WIYSNNITFI	APNTFEGFVH	LEELDLGDNR		QGLVKLHALY
Human NOGO-R3	WIYSNNITYI	HPSTFEGFVH	LEELDLGDNR		QGLVKLHALY
Human NOGO-R2	WLFSNNLSTI	YPGTFRHLOA	LEELDLGDNR		OGLERLOSLH
Consensus	WSNI		LE-LDL-DN-	-LP-TF	-Grr-
	151 LRR 4	er.	LRR 5	5	200
Human NOGO-R1	LDRCGLQELG	IDRCGLOELG PGLFRGLAAL	OYLYLODNAL OVI VI ODNNIL	QALPDDTFRD OALPDNTFRD	LGNLTHLFLH
Murine NOGO-R3	LYKCGLSALP				LUNISHLFLH
Human NOGO-R3	LYKCGLSALP			EYLQDDIFVD	LVNLSHLFLH
Human NOGO-R2	LYRCQLSSLP	GNIFRGLVBL	QYLYLQENSL	_	
Consensus	LC-LL-	F-GLL	OYLYLQ-N	L-DF-D	L-NL-HLFLH
	0				

5/6 FIG. 3, cont.

		•			
	250	RLMTLYLFAN RLMTLYLFAN RLTTLFLFUN RLTTLFLFUN RLTILYLFUN	T 300 KFRGSSSEVP KFRGSSSEVP RFRGSSSAVP RFRGSSSAVP RARVSSDVT	350 EEPLGLPKCC EELLSLPKCC SDRAARKEHH TDRAARKEHH	400 RHINDSPFGT RHINDSPFGT RNRNQISKVS RNRNQISKAG
		VHPHAFRDLG RLMTLYLFAN VHPHAFRDLG RLMTLYLFAN VHHKAFHDLH RLTTLFLFNN VHRAAFHDLR RLTTLFLFNN VHRAAFRGLS RLTILYLFNN VH-AFL- RLL-LF-N	LRRCT 300 RARPLWAWLQ KFRGSSSEVP RARSLWEWLQ KFRGSSSAVP RARSLWEWLQ RFRGSSSAVP RARPLWAWFQ RARVSSSDVT RAR-LW-WR-SSS-V-	QGCAVATGPY HPIWTGRATD EGCAVASGPF RPIQTSQLTD RNCTGPVSP, HQIKSHTLTT RNCTGPASP, HQIKSHTLTT QACP,PAAP, TRPGSRA	DSPPGNGSGP DTPPGNGSGP .KAGKNCTSH .KPGKNCTNP
	LRR 7	LLLHQNRVAH LLLHQNHVAR LLLHENQLQW LLLHENQLQW LLLHGNRLQG	LINDNPWVCDC LINGNAWDCGC LINGNPWDCGC LINGNPWDCGC LINGNPWDCGC LINANPWACDC LINANPWACDC LINANPWACDC	QGCAVATGPY EGCAVASGPF RNCTGPVSP. RNCTGPASP. QACP.PAAP.	LEPGRPASAG NALKGRVPPG LEPGRPASAG NALKGRVPPG HPHGHPPGSR SGYK HPHGPR PGHR APPADPS TLYR
FIG. 5, COIII.		AFRGLHSLDR LLLHQNRVAH VHPHAFRDLG AFRGLHSLDR LLLHQNHVAR VHPHAFRDLG IFRGLVNLDR LLLHENQLQW VHHKAFHDLH TFRGLVNLDR LLLHENQLQW VHHKAFHDLR VFRGLGSLDR LLLHGNRLQG VHRAAFRGLS -FRGL-LDR LLLH-N VHAFL-		DLKRLAANDL DLKRLAASDL DLKLLRVEDF DLKLLRAEDF DLRALREADF	
	LRR 6	•	LER8 NLSALPTEAL APLRALQYLR NLSMLPAEVL MPLRSLQYLR SLTELQGDCL APLVALEFLR SLSELQGECL APLGALEFLR SLASLPGEAL ADLPSLEFLR -LLL	301 CSLPQRLAGR CNLPQRLADR CATPELRQGQ CVSPGLRHGQ CATPPERQGR CA-P	351 QPDAADKASV QPDAADKASV PSHGASRDKG SPHGPTRSKG
		Human NOGO-R1 Murine NOGO-R3 Human NOGO-R3 Human NOGO-R2	Human NOGO-R1 Murine NOGO-R3 Human NOGO-R3 Human NOGO-R3	Human NOGO-R1 Murine NOGO-R1 Murine NOGO-R3 Human NOGO-R3	Human NOGO-R1 Murine NOGO-R1 Murine NOGO-R3 Human NOGO-R3 Consensus

6/6 FIG. 3, cont.

450 RTRSHCRLGQ RTRSHCRLGQ PIRAPSGVQQ PIRAPSGVQQ ACQAPPD	
450 LPGSAEPPLT AVRPEGSEPP GFPTSGP RRRPGCSRKN RTRSHCRLGQ LPSSAEPPLT ALRPGGSEPP GLPTTGP RRRPGCSRKN RTRSHCRLGQ .SGKELTELQ DYAPDYQHKF SFDIMPTARP KRKGKCARRT PIRAPSGVQQ .AGKQAPELP DYAPDYQHKF SFDIMPTARP KRKGKCARRT PIRAPSGVQQ .QGGDAPTED DYWGGYGGED QRGEQMCPGA ACQAPPD	AGSGGGTGD SEGSGALPSL TCSLTPLGLA LVLWTVLGPC AGSGASGTGD AEGSGALPAL ACSLAPLGLA LVLWTVLGPC ASSGASTALG APLLAWILGL AVTLR ASSASSLG ASLLAWILGL AVTLR SRGPALSA GLPS PLLCL LLLVPHHL SRGPALSA GLPS PLLCL LLLVPHHL
AVRPEGSEPP GFPTSGP RRRPGCSRKN ALRPGGSEPP GLPTTGP RRRPGCSRKN DYAPDYQHKF SFDIMPTARP KRKGKCARRT DYAPDYQHKF SFDIMPTARP KRKGKCARRT DYWGGYGGED QRGEQMCPGA	Putative GPI Signals SL TCSLTPLGLA L AL ACSLAPLGLA L G APLLAWILGL A LG ASLLAWILGL A SA GLPS PLLCL I
AVRPEGSEPP ALRPGGSEPP DYAPDYQHKF DYAPDYQHKF	Put SEGSGALPSL AEGSGALPAL ASSGTALG ASSASSLG SRGPALSA
401 LPGSAEPPLT LPSSAEPPLT SGKELTELQ AGKQAPELP	451 AGSGGGGTGD AGSGASGTGD
Human NOGO-R1 Murine NOGO-R1 Murine NOGO-R3 Human NOGO-R2 Consensus	Human NOGO-R1 Murine NOGO-R1 Murine NOGO-R3 Human NOGO-R3 Human NOGO-R2